

SEQUENCE LISTING

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<i10> Sakowicz, Roman
 Goldstein, Lawrence S. B.
 The Regents of the University of California

<120> Identification and Expression of a Novel Kinesin Motor Protein

<130> 18557C-000710US

<140> US 09/235,416

<141> 1999-01-22

<150> WO PCT/US99/01355

<151> 1999-01-22

<150> US 60/072,361

<151> 1998-01-23

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<170> PatentIn Ver. 2.0

<210> 1

<211> 784

<212> PRT

<213> Thermomyces lanuginosus

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<223> TL-gamma ATP-dependent plus end-directed microtubule motor protein

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<223> kinesin-like microtubule motor domain

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<221> DOMAIN

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<223> neck domain links motor domain to stalk domain

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<223> stalk domain, unc-104 family domain

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Arg Lys Ser Gly Lys Thr Ile Met Asp Gly Pro Lys Ala Phe Ala Phe 50 60

Asp Arg Ser Tyr Trp Ser Phe Asp Lys Asn Ala Pro Asn Tyr Ala Arg 65 70 75 80

Gln Glu Asp Leu Phe Gln Asp Leu Gly Val Pro Leu Leu Asp Asn Ala 85 90 95

Phe Lys Gly Tyr Asn Asn Cys Ile Phe Ala Tyr Gly Gln Thr Gly Ser 100 105 110

Gly Lys Ser Tyr Ser Met Met Gly Tyr Gly Lys Glu His Gly Val Ile 115 120 125

Pro Arg Ile Cys Gln Asp Met Phe Arg Arg Ile Asn Glu Leu Gln Lys 130 135 140

Asp Lys Asn Leu Thr Cys Thr Val Glu Val Ser Tyr Leu Glu Ile Tyr 145 150 150

Asn Glu Arg Val Arg Asp Leu Leu Asn Pro Ser Thr Lys Gly Asn Leu

Lys Val Arg Glu His Pro Ser Thr Gly Pro Tyr Val Glu Asp Leu Ala $180 \,\,$ $185 \,\,$ $190 \,\,$

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Gly Asn Lys Ala Arg Thr Val Ala Ala Thr Asn Met Asn Glu Thr Ser 210 215 220

Ser Arg Ser His Ala Val Phe Thr Leu Thr Leu Thr Gln Lys Trp His 225 230 235 240

Asp Glu Glu Thr Lys Met Asp Thr Glu Lys Val Ala Lys Ile Ser Leu 245 250 255

Val Asp Leu Ala Gly Ser Glu Arg Ala Thr Ser Thr Gly Ala Thr Gly 260 265 270

Ala Arg Leu Lys Glu Gly Ala Glu Ile Asn Arg Ser Leu Ser Thr Leu 275 280 285

Gly Arg Val Ile Ala Ala Leu Ala Asp Met Ser Ser Gly Lys Gln Lys 290 295 300

Lys Asn Gln Leu Val Pro Tyr Arg Asp Ser Val Leu Thr Trp Leu Leu 305 310 310 320

Lys Asp Ser Leu Gly Gly Asn Ser Met Thr Ala Met Ile Ala Ala Ile 325 330 335

Ser Pro Ala Asp Ile Asn Phe Glu Glu Thr Leu Ser Thr Leu Arg Tyr 340 345 350

Br

Ala Asp Ser Ala Lys Arg Ile Lys Asn His Ala Val Val Asn Glu Asp 360 Pro Asn Ala Arg Met Ile Arg Glu Leu Lys Glu Glu Leu Ala Gln Leu Arg Ser Lys Leu Gln Ser Ser Gly Gly Gly Gly Gly Ala Gly Gly Ser Gly Gly Pro Val Glu Glu Ser Tyr Pro Pro Asp Thr Pro Leu Glu Lys Gln Ile Val Ser Ile Gln Gln Pro Asp Ala Thr Val Lys Lys Met Ser Lys Ala Glu Ile Val Glu Gln Leu Asn Gln Ser Glu Lys Leu Tyr 435 440 Arg Asp Leu Asn Gln Thr Trp Glu Glu Lys Leu Ala Lys Thr Glu Glu 455 Ile His Lys Glu Arg Glu Ala Ala Leu Glu Glu Leu Gly Ile Ser Ile 470 475 Glu Lys Gly Phe Val Gly Pro Tyr His Ser Lys Glu Met Pro His Leu 490 Val Asn Leu Ser Asp Asp Pro Leu Leu Ala Glu Cys Leu Val Tyr Asn Ile Lys Pro Gly Gln Thr Arg Val Gly Asn Val Asn Gln Asp Thr Gln 520 Ala Glu Ile Arg Leu Asn Gly Ser Lys Ile Leu Lys Glu His Cys Thr 535 Phe Glu Asn Val Asp Asn Val Val Thr Ile Val Pro Asn Glu Lys Ala 555 Ala Val Met Val Asn Gly Val Arg Ile Asp Lys Pro Thr Arg Leu Arg Ser Gly Tyr Arg Ile Ile Leu Gly Asp Phe His Ile Phe Arg Phe Asn His Pro Glu Glu Ala Arg Ala Glu Arg Gln Glu Gln Ser Leu Leu Arg 595 His Ser Val Thr Asn Ser Gln Leu Gly Ser Pro Ala Pro Gly Arg His Asp Arg Thr Leu Ser Lys Ala Gly Ser Asp Ala Asp Gly Asp Ser Arg 630 Ser Asp Ser Pro Leu Pro His Phe Arg Gly Lys Asp Ser Asp Trp Phe 650 Tyr Ala Arg Arg Glu Ala Ala Ser Ala Ile Leu Gly Leu Asp Gln Lys 660 665



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Thr Pro Arg Ser Asp Asp Asp Gly Asp Ala Leu Phe Phe Gly Asp Lys 745

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12

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reverse primer



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